

**VERSION SHOWING CHANGES MADE**

17. (Cancel) A method comprising:

- a) providing a hybridization complex comprising
  - i) a first target sequence comprising
    - 1) a first nucleotide at a readout position; and
    - 2) a first label that uniquely identifies said first nucleotide at said readout position;
  - ii) a capture probe attached to a microsphere on a surface of a substrate, wherein said first target sequence is immobilized on said microsphere by said capture probe; and
- b) detecting said label to identify said first nucleotide at said readout position.

18.(cancel) The method according to claim 17 wherein said first target sequence comprises an adapter sequence and said adapter sequence is hybridized to said capture probe.

19.(Cancel) The method according to claim 17 further comprising:

- a) providing a second target sequence comprising a first domain and a second domain comprising a detection position;
- b) hybridizing a first ligation probe to said first domain and a second ligation probe to said second domain of said second target sequence wherein if said second

ligation probe comprises a nucleotide that is perfectly complementary to said nucleotide at said detection position a ligation structure is formed;

c) ligating said ligation structure to form said first target sequence, wherein said readout position is perfectly complementary to said detection position.

20. (cancel) The method according to claim 19, wherein said first ligation probe comprises an adapter sequence and said second ligation probe comprises said first label.

21. (cancel) The method according to claim 17 further comprising:

- a) providing a second target sequence comprising said detection position;
- b) hybridizing an extension primer to said second target sequence adjacent to said detection position;
- c) adding a polymerase enzyme and at least a first dNTP comprising a covalently attached detectable label under conditions whereby if said first dNTP basepairs with the nucleotide at said detection position, said extension primer is extended by said enzyme to incorporate said label into said extension primer to form said first target sequence.

22. (cancel) The method according to claim 21, further comprising adding a second dNTP, wherein said first and second dNTPs comprise first and second labels, respectively.

23.(Amended) The method according to claim 42, wherein at least said [first] detectable label comprises a fluorophore.

24.(Amended) The method according to claim 42, wherein at least said [first] detectable label comprises biotin.

25.(Amended) The method according to claim 42, wherein at least said [first] detectable label comprises imine-biotin.

26.(Amended) The method according to claim 42, wherein said [at least] said [first] dNTP comprises a functional group for addition of a fluorophore.

27. (cancel) The method according to claim 17 further comprising:

a) providing a second target sequence comprising 5' to 3':

i) a first target domain comprising an overlap domain comprising at least a nucleotide in the detection position; and

ii) a second target domain contiguous with said detection position;

b) hybridizing:

i) a first probe to said first target domain; and

ii) a second probe to said second target domain, wherein said second probe comprises:

1) a detection sequence that does not hybridize with said target sequence; and

2) a detectable label;

wherein if said second probe comprises a nucleotide that is perfectly complementary to said detection position a cleavage structure is formed; and

c) contacting said cleavage structure with a cleavage enzyme to cleave said detection sequence to form said first target sequence.

28.(cancel) The method according to claim 19, 21 or 27, wherein said first target sequence comprises an adapter sequence and said adapter sequence is hybridized to said capture probe.

29.(Amended) The method according to claim [17, 19, 21 or 27] 42, wherein said substrate is a fiber optic bundle.

30. The method according to claim [17, 19, 21 or 27] 42 wherein said substrate is selected from the group consisting of glass and plastic.

31.(Amended) The method according to claim [17, 18, 19, 21 or 27] 42, wherein said [first] detectable label is a fluorophore.

32.(cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

a) providing a hybridization complex comprising:

i) a target sequence; and

ii) at least a first probe, wherein said first probe is hybridized to said target sequence adjacent to said detection position;

b) adding a composition comprising:

i) a nucleotide that hybridizes with the nucleotide at said detection position; and

ii) an enzyme, wherein said enzyme alters said first probe when said nucleotide hybridizes with said nucleotide at said detection position to form an altered probe, wherein said altered probe comprises a label specific to said nucleotide;

c) forming an assay complex by hybridizing said altered probe with a capture probe covalently attached to a microsphere on a surface of a substrate; and

d) determining the nucleotide at said detection position by detecting said label.

33. (cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

a) providing a hybridization complex comprising a target sequence, at least a first probe hybridized with said target sequence adjacent to said detection position, and a capture probe covalently attached to a microsphere on a surface of a substrate;

b) adding a composition comprising a nucleotide that hybridizes with said detection position and an enzyme, wherein said enzyme alters said first probe

when said nucleotide hybridizes with said detection position to form an altered probe, wherein said altered probe comprises a label that uniquely identifies said nucleotide hybridized with said detection position; and  
d) determining the nucleotide at said detection position by detecting said label.

34. (cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

- a) providing a hybridization complex comprising a target sequence and a capture probe covalently attached to a microsphere on a surface of a substrate, wherein said capture probe hybridizes to said target sequence;
- b) adding a composition comprising a nucleotide that hybridizes with said detection position and an enzyme, wherein said enzyme alters said capture probe when said nucleotide hybridizes with said detection position to form an altered capture probe, wherein said altered capture probe comprises a label; and
- d) determining the nucleotide at said detection position by detecting said label.

35. (cancel) The method according to claim 32, 33 or 34, wherein said label is a fluorophore.

36. (cancel) The method according to claim 32, 33 or 34, wherein said nucleotide is a first dNTP comprising a first label and said enzyme is a polymerase, whereby when said first dNTP basepairs with the nucleotide at said detection position, said first probe is extended by said enzyme to incorporate said first label into said first probe.

37.(cancel) The method according to claim 32, 33 or 34, wherein said composition comprises a second probe comprising said nucleotide wherein said second probe hybridizes with said target sequence, said nucleotide basepairs with said detection position and said enzyme is a ligase, whereby when said nucleotide basepairs with said nucleotide at said detection position, a ligation structure is formed and said ligase ligates said ligation structure.

38. (cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

a) providing a hybridization complex comprising said target sequence hybridized with a capture probe, wherein said capture probe is covalently attached to a microsphere on a surface of a substrate; and

b) contacting said hybridization complex with a plurality of detection probes each comprising:

i) a unique nucleotide at a readout position; and

ii) a unique detectable label; and

c) detecting a signal from at least one of said detectable labels to identify the nucleotide at the detection position.

39. (cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

a) providing a hybridization complex comprising said target sequence, wherein said target sequence comprises a first target domain directly 5' adjacent to said detection position, a

capture probe covalently attached to a microsphere on a surface of a substrate, and an extension primer hybridized to said first target domain of said target sequence, wherein said capture probe hybridizes with either said extension primer or said target sequence;

b) contacting said hybridization complex with:

i) a polymerase enzyme;

ii) a plurality of dNTPs each comprising a covalently attached detectable label;

under conditions whereby if one of said dNTPs basepairs with the nucleotide at said detection position, said extension primer is extended by said enzyme to incorporate said label; and

c) identifying the nucleotide at said detection position.

40. (cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising a first target domain comprising said detection position and a second target domain adjacent to said detection position, said method comprising:

a) hybridizing a first ligation probe to said first target domain;

b) hybridizing a second ligation probe to said second target domain, wherein if said second ligation probe comprises a nucleotide that is perfectly complementary to said detection position a ligation structure is formed;

c) ligating said first and said second ligation probes to form a ligated probe;

d) forming a complex with said ligated probe, a capture probe covalently attached to a microsphere on a surface of a substrate, and at least one label;



- e) detecting the presence or absence of said label as an indication of the formation of said ligation structure; and
- f) identifying the nucleotide at said detection position.

41. (cancel) A method of determining the identification of a nucleotide at a detection position in a target sequence wherein said target sequence comprises 5' to 3':

- a) a first target domain comprising an overlap domain comprising at least a nucleotide in the detection position; and

- b) a second target domain contiguous with said detection position;

said method comprising:

- i) providing a hybridization complex, wherein said hybridization complex comprises:

- 1) a first probe hybridized to said first target domain; and
- 2) a second probe hybridized to said second target domain,

wherein said second probe comprises:

- i) a detection sequence that does not hybridize with said target sequence; and
- ii) a detectable label;

wherein if said second probe comprises a nucleotide that is perfectly complementary to said detection position a cleavage structure is formed;

- ii) contacting said hybridization complex with a cleavage enzyme that will cleave said detection sequence;

iii) forming a complex with said detection sequence, a capture probe covalently attached to a microsphere on a surface of a substrate, and at least one label; and  
iv) detecting the presence or absence of said label as an indication of the formation of said cleavage structure, whereby the nucleotide at said detection is identified.

42. (NEW) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

a) providing a hybridization complex comprising

i) a first target sequence comprising

1) a first nucleotide at a detection position; and

2) a first target domain directly 5' adjacent to said detection position;

3) a second target domain 3' adjacent to said detection position;

ii) a first ligation probe hybridized to said first target domain;

iii) a second ligation probe hybridized to said second target domain;

b) contacting said hybridization complex with:

i) an extension enzyme;

ii) at least one dNTP;

such that if the base of said dNTP is perfectly complementary to the base of said detection position, said first ligation probe is extended to form a ligation structure;

c) contacting said ligation structure with a ligase to ligate said first extended

ligation probe and said second ligation probe to form a ligation product; and

d) detecting the presence of said ligation product to identify the nucleotide at said detection position, said detecting comprising providing a substrate with a surface comprising discrete sites, further comprising a population of microspheres comprising at least a first and a second subpopulation, wherein each subpopulation comprises a capture probe, wherein said capture probe hybridizes to a sequence contained within said ligation product.

- 43.(NEW) The method according to claim 42 wherein one of said ligation probe comprises an adapter sequence that hybridizes to said capture probe.
- 44.(NEW) The method according to claim 42 wherein said dNTP comprises a detectable label.
45. (NEW) The method according to claim 42 wherein said capture probe attached to a microsphere on a surface of said substrate serves as said first ligation probe.
- 47.(NEW) The method according to claim 42, wherein said capture probe is a nucleic acid.
- 48.(NEW) The method according to claim 42, wherein said capture probe is a protein.
- 49.(NEW) The method according to claim 42, wherein said discrete sites are wells.

50.(NEW) The method according to claim 42, wherein said microspheres are randomly distributed on said substrate.

## APPENDIX A

### Pending Claims

23. The method according to claim 42, wherein said detectable label comprises a fluorophore.
24. The method according to claim 42, wherein said detectable label comprises biotin.
25. The method according to claim 42, wherein said detectable label comprises imine-biotin.
26. The method according to claim 42, wherein said dNTP comprises a functional group for addition of a fluorophore.
29. The method according to claim 42, wherein said substrate is a fiber optic bundle.
30. The method according to claim 42, wherein said substrate is selected from the group consisting of glass and plastic.
31. The method according to claim 42, wherein said detectable label is a fluorophore.

42. (NEW) A method of determining the identification of a nucleotide at a detection position in a target sequence comprising:

a) providing a hybridization complex comprising

i) a first target sequence comprising

1) a first nucleotide at a detection position; and

2) a first target domain directly 5' adjacent to said detection position;

3) a second target domain 3' adjacent to said detection position;

ii) a first ligation probe hybridized to said first target domain;

iii) a second ligation probe hybridized to said second target domain;

b) contacting said hybridization complex with:

i) an extension enzyme;

ii) at least one dNTP;

such that if the base of said dNTP is perfectly complementary to the base of said detection position, said first ligation probe is extended to form a ligation structure;

c) contacting said ligation structure with a ligase to ligate said first extended ligation probe and said second ligation probe to form a ligation product; and

d) detecting the presence of said ligation product to identify the nucleotide at said detection position, said detecting comprising providing a substrate with a surface comprising discrete sites, further comprising a population of microspheres comprising at least a first and a second subpopulation, wherein each subpopulation comprises a capture probe, wherein said capture probe hybridizes to a sequence contained within said ligation product.

- 43.(NEW) The method according to claim 42 wherein one of said ligation probe comprises an adapter sequence that hybridizes to said capture probe.
- 44.(NEW) The method according to claim 42 wherein said dNTP comprises a detectable label.
45. (NEW) The method according to claim 42 wherein said capture probe attached to a microsphere on a surface of said substrate serves as said first ligation probe.
- 47.(NEW) The method according to claim 42, wherein said capture probe is a nucleic acid.
- 48.(NEW) The method according to claim 42, wherein said capture probe is a protein.
- 49.(NEW) The method according to claim 42, wherein said discrete sites are wells.
- 50.(NEW) The method according to claim 42, wherein said microspheres are randomly distributed on said substrate.